

Supporting Information

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SI Materials and Methods

Antibodies, Reagents, and Medium. The following mAbs were purchased from BD Bioscience (San Jose, CA) or eBioscience (San Diego, CA): functional grade purified mAbs for CD3 ϵ (145–2C11) and CD28 (37.51); biotin-, FITC-, PE-, PerCP-Cy5- or allophycocyanin-labeled mAbs for human CD2 (hCD2) (RPA-2.10), CD4 (RM4–5), CD25 (PC61 or 7D4), CD45.1 (A20), CD45.2 (104), CTLA-4 (UC10–4F10–11), GITR (DTA-1), TCR- β (H57–597), V α 2 (B20.1), V β 3 (KJ25), V β 4 (KT4), V β 5.1/5.2 (MR9–4), V β 6 (RR4–7), V β 8.1/8.2 (MR5–2), V β 14 (14–2), IFN- γ (XMG1.2), IL-2 (JES6–5H4), IL-4 (11B11), and IL-17A (eBio17B7). Anti-TGF- β 1/2/3 mAbs (1D11), recombinant mouse (rm) TGF β RII-Fc chimeric proteins, rmIL-2 and recombinant human TGF- β 1 were purchased from R&D systems (Minneapolis, MN), and rmIL-4, rmIL-6, rmIL-12 were obtained from PeproTech (London, UK). T cells were cultured in RPMI10 medium; RPMI-1640 supplemented with 10% fetal bovine serum, 10 mmol/L HEPES, 1 mmol/L sodium pyruvate, 50 μ mol/L 2-ME, 100 U/ml penicillin, and 100 μ g/ml streptomycin.

Cell Sorting. To purify peripheral CD4 $^{+}$ T cell subsets, pooled spleen and lymph node (LN) cells were first depleted of B220 $^{+}$, CD8 $^{+}$, CD11b $^{+}$, Gr1 $^{+}$, and adherent cells by panning, stained with fluorescent mAbs, and sorted on a FACS Aria cell sorter. To sort CD4 $^{+}$ Foxp3 $^{+}$ CD25 high and CD25 $^{-}$ T cells, Foxp3 $^{hCD2^{+}}$ T cells were first enriched using the MACS system. In brief, enriched CD4 $^{+}$ T cells were stained with FITC-anti-CD4, PE-anti-hCD2, and biotin-anti-CD25 followed by streptavidin-PE-Cy5, labeled with anti-PE microbeads, and passed through LS columns (Miltenyi Biotec) to obtain hCD2 $^{+}$ cells, which were further FACS sorted into CD4 $^{+}$ Foxp3 $^{hCD2^{+}}$ CD25 high and CD25 $^{-}$ T cells.

CFSE Labeling. Sorted T cells were incubated with 5 μ mol/L (for in vitro suppression assay) or 10 μ mol/L (for adoptive transfer) CFSE in PBS containing 0.1% bovine serum albumin for 10 minutes at 37 $^{\circ}$ C. The reaction was stopped by adding $\times 10$ volume of RPMI10 medium, and cells were washed four times with RPMI10 medium.

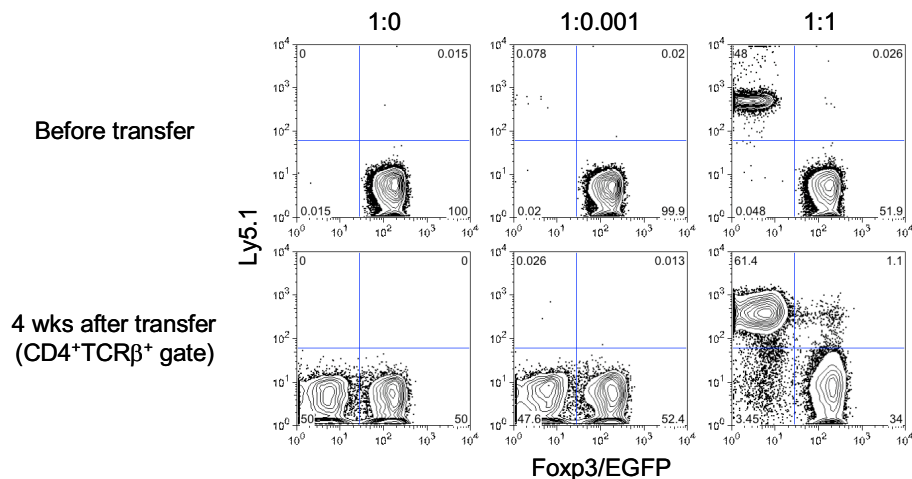
Initial Foxp3⁺ : Foxp3⁻ ratio

Fig. S1. The generation of Foxp3⁻ T cells from Foxp3⁺ T cells is not caused by outgrowth of Foxp3⁻ T cells contaminating the injected donor cells, but reflects loss of Foxp3 expression in Foxp3⁺ T cells. CD4⁺EGFP⁺ T cells sorted from Foxp3^{EGFP} Ly5.2 mice were mixed with CD4⁺EGFP⁻ T cells sorted from Foxp3^{EGFP} Ly5.1 mice at the indicated ratio and adoptively transferred into RAG2^{-/-} host mice as described in Fig. 1A. Upper panels depict Ly5.1 and EGFP expression profiles of the injected populations. Lower panels show Ly5.1 and EGFP expression on LN CD4⁺TCRβ⁺ cells analyzed 4 weeks after transfer.

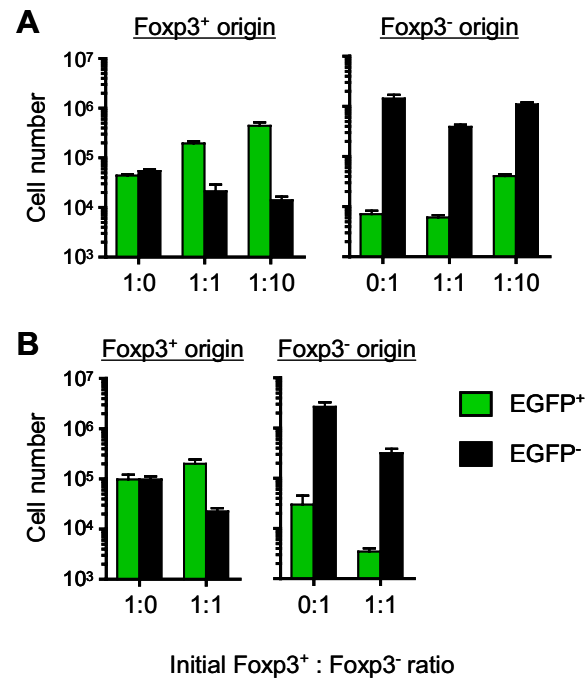


Fig. S2. Foxp3 down-regulation in RAG2^{-/-} and CD3 ϵ ^{-/-} recipients. Sorted CD4⁺Foxp3/EGFP⁺ and EGFP⁻ T cells were injected into RAG2^{-/-} (A) or CD3 ϵ ^{-/-} mice (B) as described in Fig. 1A. Four weeks after transfer, LN and spleen (not depicted) cells were counted and stained for CD4, TCR β , and Ly5.1. Absolute numbers of Foxp3⁺ (left) or Foxp3⁻ (right) donor-derived T cells with the indicated EGFP phenotype are shown ($n = 4-5$ per group).

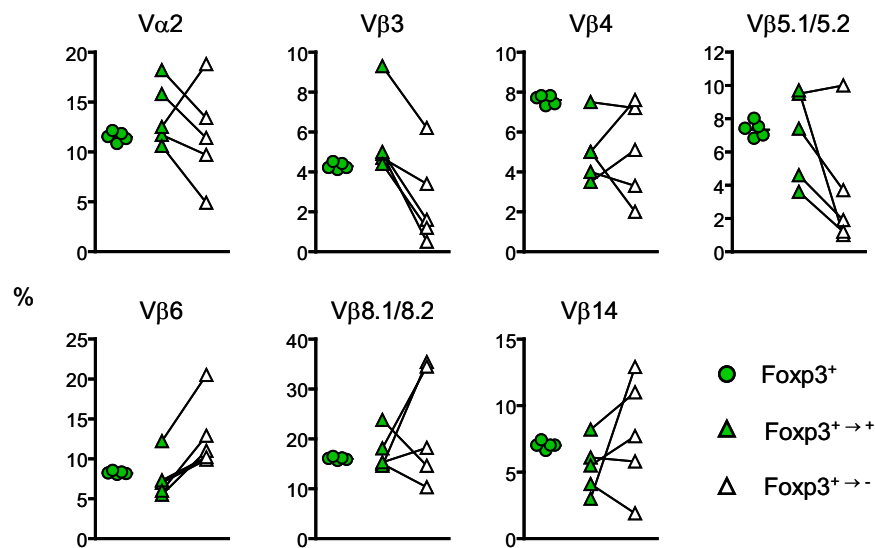


Fig. S3. TCR repertoires of Foxp3-maintaining and Foxp3-down-regulated T cells. CD4⁺EGFP⁺ T cells sorted from Foxp3^{EGFP} Ly5.1 mice were injected into CD3ε^{-/-} host mice (1 × 10⁵). Four weeks after transfer, LN and spleen cells from individual host mice were pooled, enriched for donor T cells by depleting B220⁺, CD8⁺, CD11b⁺, and Gr1⁺ cells by panning, and stained for Ly5.1 and the indicated TCR Vα or Vβ chains. Frequencies of cells expressing each Vα/Vβ chain among the Foxp3⁺ and Foxp3⁺ cells as well as CD4⁺EGFP⁺ T cells from Foxp3^{EGFP} mice were determined (n = 5 per group). Lines connect Foxp3⁺ and Foxp3⁺ cells from each individual host.

